

logo

NTDB id 378 SMU RS02690 WP 002263569.1	MLCQNCCKINEA.....TI	13
NTDB id 241 SP RS04015 WP 000882523.1	MLCQNCCKINDS.....TI	13
NTDB id 281 KZH43 RS03550 WP 000882517.1	MLCQNCCKINDS.....TI	13
NTDB id 245 SPD RS03860 WP 000882517.1	MLCQNCCKINDS.....TI	13
NTDB id 243 SPR RS03635 WP 000882517.1	MLCQNCCKINDS.....TI	13
NTDB id 85 BSU 00860 NP 387967.1	..MMFGRFTERAQKVLALAQEEALRLGHNNIGTEHILLGLVREGEGIAAKALQALGLGSEKIQKEVESLIG.....RGQEMS	75
NTDB id 610 V4T04 RS10165 WP 012897346.1	MKFENIKYTPPTLDRILEKAEYEAHQYQYGTIESAHLAAMATTSGSIAYSLLAGMNVDSDDLIDL.....EDLSSHVKVKR	77
NTDB id 287 SP RS11210 WP 001109712.1	MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNHSYSVAGATLNDYPYEMDRLE....EVALELTETDYSQDET	75
NTDB id 290 KZH43 RS10025 WP 001109677.1	MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNHSYSVAGATLNDYPYEMDRLE....EVALELTETDYSQDET	75
NTDB id 289 SPD RS10700 WP 001109677.1	MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNHSYSVAGATLNDYPYEMDRLE....EVALELTETDYSQDET	75
NTDB id 377 SMU RS09275 WP 002262344.1	MTDYSLKMQEVFRLAQFEAARFESSYLESWHVLLAMVEIDSSVAGLSFAEFEADVRFEDYQAAAILAIGKKPKSSSN..DI	79
NTDB id 329 STU RS10020 WP 011225298.1	MTIYSRKMQAIFHRAQLEAERFESPFLLETWHVLLAMVEVPGSVAYLTFTDFEDRIHSEEIETAAVLAMEKRPKDLSESDII	81
NTDB id 297 STER RS00545 WP 011680614.1	MTIYSRKMQAIFHRAQLEAERFESPFLLETWHVLLAMVEVTGSAVAYLTFTDFEDRIHSEEIETAAVLAMEKRPKDLSESDII	81
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NTDB id 378 SMU RS02690 WP 002263569.1	HLIYNSNRQLQKQIDLA.....QNCYQIMKTDPENGLNHLISQ.....QNSSINPFFDDFFGDLNNFRAFNNQDL	78
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NTDB id 243 SPR RS03635 WP 000882517.1	HLIYTNLNGKQKQIDLC.....QNCYKIKTDPNNSLFKGMTD.....LNNRDFDP.FGDFFNDLNNFRPSSN..T	75
NTDB id 85 BSU 00860 NP 387967.1	QTIHYTPRAKKVIELSMDEARKLGHSYVGTEHILLGLIRE.G.EGVAARVLNNLGVSLNKARQQVLQL.....LGSNET..	147
NTDB id 610 V4T04 RS10165 WP 012897346.1	STLRFSPRAEEVMTAASFLAIHNNSEAVGTEHLLYALLQV.E.DGFGQLLKL.....QKINIVSLRKELEKRTGLKVPES	151
NTDB id 287 SP RS11210 WP 001109712.1	TELPFSRRLQVLFDAAEYVASVVHAKVLGTEHVLYAILHD.S.NALATRILERAGFSYEDKKDQVKIAALRRNLEERAGWTREDL	158
NTDB id 290 KZH43 RS10025 WP 001109677.1	TELPFSRRLQILFDAAEYVASVVHAKVLGTEHVLYAILHD.G.NALATRILERAGFSYEDKKDQVKIAALRRNLEERAGWTREDL	158
NTDB id 289 SPD RS10700 WP 001109677.1	TELPFSRRLQILFDAAEYVASVVHAKVLGTEHVLYAILHD.G.NALATRILERAGFSYEDKKDQVKIAALRRNLEERAGWTREDL	158
NTDB id 377 SMU RS09275 WP 002262344.1	MLLEQSHALKRITLAEAAIASVQTHAKEVGTEHVLFAMLLN.P.NLLATRILELVGFHAKDDGESIRLLDLRKVIERYACFSKEDI	162
NTDB id 329 STU RS10020 WP 011225298.1	DLRAQSPALEAMLQEAQGIASVTGAVEVGSEHVLMAFLLH.K.DLMVCRILLEVAGFQYKDDSDKPRIIDLRRSLERNAGLSKQDL	164
NTDB id 297 STER RS00545 WP 011680614.1	DLRAQSPALEAMLQEAQEIASVTGAVEVGSEHVLMAFLLH.K.DLMVCRILLEVAGFQYKDDSDKPRIIDLRRSLERNAGLSKQDL	164
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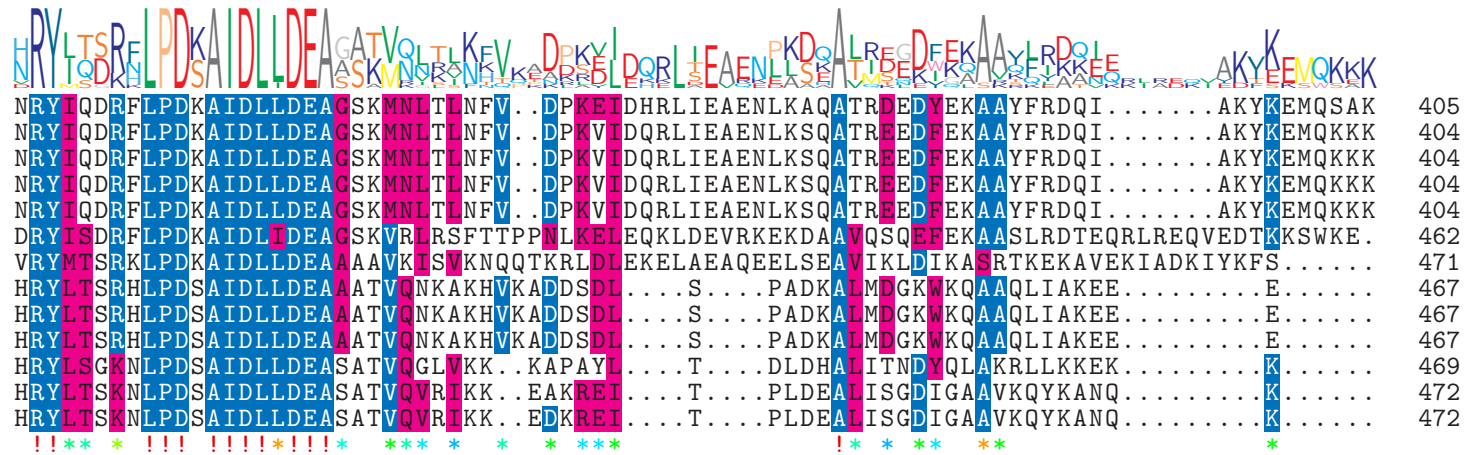
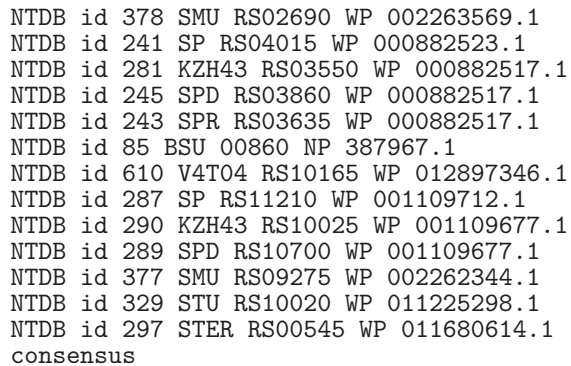
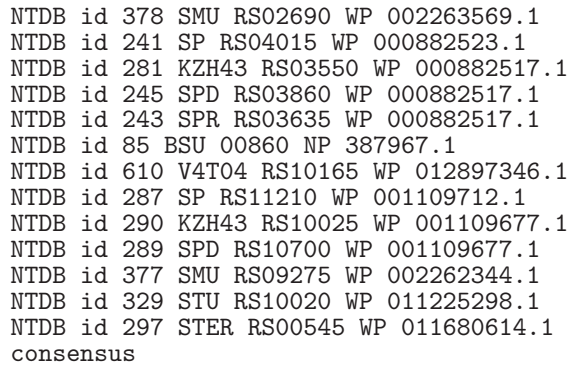
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NTDB id 281 KZH43 RS03550 WP 000882517.1	PP.IPPTQSGGGYGGNGGYGSQNRGSAQTPPPSQEKGLEEFGINVTEIARRGIDPVIGRDDEIIRVIEILNRRRTKNNPVLIGE	159
NTDB id 245 SPD RS03860 WP 000882517.1	PP.IPPTQSGGGYGGNGGYGSQNRGSAQTPPPSQEKGLEEFGINVTEIARRGIDPVIGRDDEIIRVIEILNRRRTKNNPVLIGE	159
NTDB id 243 SPR RS03635 WP 000882517.1	PP.IPPTQSGGGYGGNGGYGSQNRGSAQTPPPSQEKGLEEFGINVTEIARRGIDPVIGRDDEIIRVIEILNRRRTKNNPVLIGE	159
NTDB id 85 BSU 00860 NP 387967.1	GSSAAGTNSNANTPTLDSLARDLTAIKEDSLDPVIGRSKEIQRVIEVLSRRTKNNPVLIGE	209
NTDB id 610 V4T04 RS10165 WP 012897346.1	KKAVTPMSKRKM.....AKG.....VAENSTPTLDSVSSDLTEEARLGLKLDPMIGRAEIDRLIHILSRRTKNNPVLIGE	222
NTDB id 287 SP RS11210 WP 001109712.1	KA.LRQRHRTV.....ADKQNSMANMMGMPQTPSGGLEDYTHDLTEQARSQKLEPVIGRDKIEISRMIIQLSRKTNNPVLVGD	235
NTDB id 290 KZH43 RS10025 WP 001109677.1	KA.LRQRHRTV.....ADKQNSMANMMGMPQTPSGGLEDYTHDLTEQARSQKLEPVIGRDKIEISRMIIQLSRKTNNPVLVGD	235
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NTDB id 377 SMU RS09275 WP 002262344.1	KA.IFEMRPKK.....VKNSSSFSDLMKPP.SVTGELADFTRDLTELARQGRLEPVIGRDEEISRMVQILSRKTNNPVLVGD	239
NTDB id 329 STU RS10020 WP 011225298.1	KA.IHDLRKPKK.....SKASANFANMMQPPQSSTGELADYTKDLTALAESGNLDPVIGRDEEISRMIIQLSRKTNNPVLVGD	242
NTDB id 297 STER RS00545 WP 011680614.1	KA.IHDLRKPKK.....SKASANFANMMQPPQSSTGELADYTKDLTALAESGNLDPVIGRDEEISRMIIQLSRKTNNPVLVGD	242
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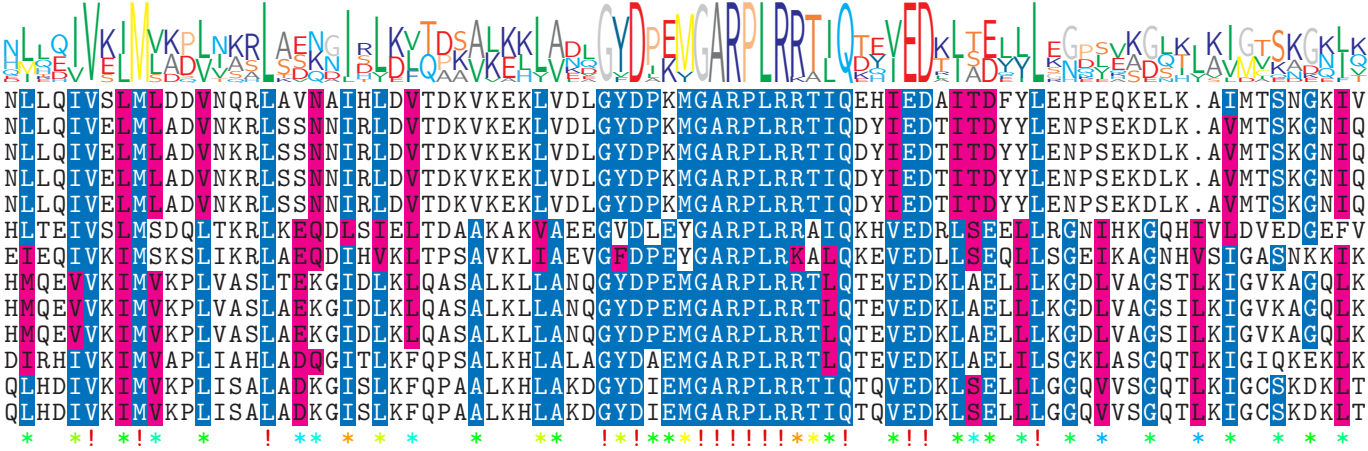
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consensus



737
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logo

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NTDB id 329 STU RS10020 WP 011225298.1
NTDB id 297 STER RS00545 WP 011680614.1
consensus



non conserved
similar
≥ 50% conserved